RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: ___10/538, 291___

Source: ___JEWP___

Date Processed by STIC: ___06/29/2006___

ENTERED

RAW SEQUENCE LISTING   DATE: 06/29/2006
PATENT APPLICATION: **US/10/538,291**   TIME: 13:48:04

Input Set : **A:\PTO.DA.txt**
Output Set: **N:\CRF4\06292006\J538291.raw**

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      3 <110> APPLICANT: SHOWA DENKO K.K.
W-->  4 <120> TITLE OF INVENTION: Method for screening E. coli strains which exhibit enhanced
W-->  5       express of a foreign gene, such E. coli strains and method for
W-->  6       producing protein or other compounds using the same
W-->  7 <130> FILE REFERENCE: SDF4812PCT
C-->  8 <140> CURRENT APPLICATION NUMBER: US/10/538,291
C-->  8 <141> CURRENT FILING DATE: 2005-06-10
      8 <150> PRIOR APPLICATION NUMBER: JP 2002-360564
      9 <151> PRIOR FILING DATE: 2002-12-12
W--> 10 <160> NUMBER OF SEQ ID: 2
     11 <170> SOFTWARE: PatentIn Ver. 2.1
     13 <210> SEQ ID NO: 1
     14 <211> LENGTH: 18
     15 <212> TYPE: DNA
     16 <213> ORGANISM: Lithospermum erythrorhizon
     18 <400> SEQUENCE: 1
     19 taagcgtaat tagctgag                                        18
     21 <210> SEQ ID NO: 2
     22 <211> LENGTH: 18
     23 <212> TYPE: DNA
     24 <213> ORGANISM: Lithospermum erythrorhizon
     26 <400> SEQUENCE: 2
     27 agatcgaaga tctaaaca                                        18
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VERIFICATION SUMMARY　　　　　　　　DATE: 06/29/2006
PATENT APPLICATION: **US/10/538,291**　　　TIME: 13:48:05

Input Set : **A:\PTO.DA.txt**
Output Set: **N:\CRF4\06292006\J538291.raw**

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L:4 M:283 W: Missing Blank Line separator, <120> field identifier
L:7 M:283 W: Missing Blank Line separator, <130> field identifier
L:8 M:270 C: Current Application Number differs, Replaced Current Application No
L:8 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:10 M:283 W: Missing Blank Line separator, <160> field identifier
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CRF Errors Edited by the STIC Systems Branch

Serial Number: __10/538, 291__   CRF Edit Date: __06/29/2006__
                                    Edited by: __DA__

____ Realigned nucleic acid/amino acid numbers/text in cases where the sequence
text "wrapped" to the next line

____ Corrected the SEQ ID NO. Sequence numbers edited were:

_____

____ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID
NO's edited:

_____

__/__ Deleted: ___ invalid beginning/end-of-file text ; ___ page numbers

____ Inserted mandatory headings/numeric identifiers, specifically:

_____

____ Moved responses to same line as heading/numeric identifier, specifically:

_____

____ Other:

_____
_____
_____

Revised 09/09/2003

Raw Sequence Listing before editing (for reference only)

IFWP

RAW SEQUENCE LISTING   DATE: 06/29/2006
PATENT APPLICATION: US/10/538,291   TIME: 10:21:32

        Input Set : A:\PTO.RJ.txt
        Output Set: N:\CRF4\06292006\J538291.raw

          3 <110> APPLICANT: SHOWA DENKO K.K.
W-->   4 <120> TITLE OF INVENTION: Method for screening E. coli strains which exhibit enhanced
W-->   5   express of a foreign gene, such E. coli strains and method for
W-->   6   producing protein or other compounds using the same
W-->   7 <130> FILE REFERENCE: SDF4812PCT
C-->   8 <140> CURRENT APPLICATION NUMBER: US/10/538,291
C-->   8 <141> CURRENT FILING DATE: 2005-06-10
          8 <150> PRIOR APPLICATION NUMBER: JP 2002-360564
          9 <151> PRIOR FILING DATE: 2002-12-12
W--> 10 <160> NUMBER OF SEQ ID: 2
         11 <170> SOFTWARE: PatentIn Ver. 2.1

Does Not Comply
Corrected Diskette Needed

ERRORED SEQUENCES

         21 <210> SEQ ID NO: 2
         22 <211> LENGTH: 18
         23 <212> TYPE: DNA
         24 <213> ORGANISM: Lithospermum erythrorhizon
         26 <400> SEQUENCE: 2
         27 agatcgaaga tctaaaca   18
E--> 34 1
E--> 36 1/2

Deleted

VERIFICATION SUMMARY   DATE: 06/29/2006
PATENT APPLICATION: US/10/538,291   TIME: 10:21:33

Input Set : **A:\PTO.RJ.txt**
Output Set: **N:\CRF4\06292006\J538291.raw**

L:4 M:283 W: Missing Blank Line separator, <120> field identifier
L:7 M:283 W: Missing Blank Line separator, <130> field identifier
L:8 M:270 C: Current Application Number differs, Replaced Current Application No
L:8 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:10 M:283 W: Missing Blank Line separator, <160> field identifier
L:34 M:254 E: No. of Bases conflict, this line has no nucleotides.
M:254 Repeated in SeqNo=2
L:36 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:36 M:252 E: No. of Seq. differs, <211> LENGTH:Input:18 Found:19 SEQ:2